1/22

## PCT/US2005/001630 IAP6 Rac'd PCT/PTO 24 JUL 2006

## SEQUENCE LISTING

<110>	Zimmerman	ı, et al	•							
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Tyr Leu Lys Lys Ala Phe Leu Leu Val Gln Asp Ile Met Glu Asp Thr

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Gln Glu Leu Ser Leu Arg Leu Lys Ser Cys Phe Thr Lys Asp Tyr Glu 115

Glu His Asp Lys Ala Cys Val Arg Thr Phe Tyr Glu Thr Pro Leu Gln

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atc acc gag gag gtg tcg gag tac tgt agc cac atg att ggg agt gga 327 Ile Thr Glu Glu Val Ser Glu Tyr Cys Ser His Met Ile Gly Ser Gly 35 40

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		aag Lys						615	5
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Tyr Leu Lys Lys Ala Phe Leu Leu Val Gln Asp Ile Met Glu Asp Thr

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Gln Lys Thr Asp His Pro Ser Ala Leu Leu Arg Asp Pro Pro Glu Pro 390

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Ser Thr Leu Ser Ala Gln Pro Gln Leu Ser Arg Ser His Ser Ser Gly

Ser Val Leu Pro Leu Gly Glu Leu Glu Gly Arg Arg Ser Thr Arg Asp

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Ala Arg Pro Leu Pro Arg Phe Asn Ser Val Pro Leu Thr Asp Thr Gly 470

His Glu Arg Gln Ser Glu Gly Ser Ser Ser Pro Gln Leu Gln Glu Ser

Val Phe His Leu Leu Val Pro Ser Val Ile Leu Val Leu Leu Ala Val 500 505

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Thr Glu Leu Ser Pro Ser Arg Pro Gly Gly Gly Ser Met Gln Thr Glu

Pro Ala Arg Pro Ser Asn Phe Leu Ser Ala Ser Ser Pro Leu Pro Ala

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	ctc Leu															778
	atc Ile															826
	ctg Leu 180															874
gtg Val 195	cag Gln	aaa Lys	gtc Val	atc Ile	cca Pro 200	G1y 999	ccc Pro	cca Pro	gcc Ala	ttg Leu 205	aca Thr	ctg Leu	gtg Val	cct Pro	gca Ala 210	922
gag Glu	ctg Leu	gtg Val	cgg Arg	att Ile 215	cga Arg	gjå aaa	gag Glu	gct Ala	gcc Ala 220	cag Gln	atc Ile	gtg Val	tgc Cys	tca Ser 225	gcc Ala	970
agc Ser	agc Ser	gtt Val	gat Asp 230	gtt Val	aac Asn	ttt Phe	gat Asp	gtc Val 235	ttc Phe	ctc Leu	caa Gln	cac His	aac Asn 240	aac Asn	acc Thr	1018
	ctc Leu															1066
aaa Lys	gtc Val 260	ctg Leu	acc Thr	ctc Leu	aac Asn	ctc Leu 265	gat Asp	caa Gln	gta Val	gat Asp	ttc Phe 270	caa Gln	cat His	gcc Ala	ggc Gly	1114
aac Asn 275	tac Tyr	tcc Ser	tgc Cys	gtg Val	gcc Ala 280	agc Ser	aac Asn	gtg Val	cag Gln	ggc Gly 285	aag Lys	cac His	tcc Ser	acc Thr	tcc Ser 290	1162
atg Met	ttc Phe	ttc Phe	cgg Arg	gtg Val 295	gta Val	gag Glu	agt Ser	gcc Ala	tac Tyr 300	ttg Leu	aac Asn	ttg Leu	agc Ser	tct Ser 305	gag Glu	1210
cag Gln	aac Asn	ctc Leu	atc Ile 310	cag Gln	gag Glu	gtg Val	acc Thr	gtg Val 315	gjå aaa	gag Glu	gjå aaa	ctc Leu	aac Asn 320	ctc Leu	aaa Lys	1258
gtc Val	atg Met	gtg Val 325	gag Glu	gcc Ala	tac Tyr	cca Pro	ggc 330	ctg Leu	caa Gln	ggt Gly	ttt Phe	aac Asn 335	tgg Trp	acc Thr	tac Tyr	1306
ctg Leu	gga Gly 340	ccc Pro	ttt Phe	tct Ser	gac Asp	cac His 345	cag Gln	cct Pro	gag Glu	ccc Pro	aag Lys 350	ctt Leu	gct Ala	aat Asn	gct Ala	1354
acc Thr 355	acc Thr	aag Lys	gac Asp	aca Thr	tac Tyr 360	agg Arg	cac His	acc Thr	ttc Phe	acc Thr 365	ctc Leu	tct Ser	ctg Leu	ccc Pro	cgc Arg 370	1402
ctg Leu	aag Lys	ccc Pro	tct Ser	gag Glu	gct Ala	ggc Gly	cgc Arg	tac Tyr	tcc Ser	ttc Phe	ctg Leu	gcc Ala	aga Arg	aac Asn	cca Pro	1450

				375					380					385			
			_	_	_	acg Thr						_					1498
						aca Thr											1546
						ccc Pro 425											1594
						tgt Cys											1642
						gtc Val											1690
						act Thr											1738
tac Tyr	gag Glu	tgc Cys 485	agg Arg	gcc Ala	cac His	aac Asn	agc Ser 490	gtg Val	gjå aaa	agt Ser	ggc Gly	tcc Ser 495	tgg Trp	gcc Ala	ttc Phe		1786
ata Ile	Pro 500	atc Ile	tct Ser	gca Ala	gga Gly	gcc Ala 505	cac His	acg Thr	cat His	ccc Pro	ccg Pro 510	gat Asp	gag Glu	ttc Phe	ctc Leu		1834
ttc Phe 515	aca Thr	cca Pro	gtg Val	gtg Val	gtc Val 520	gcc Ala	tgc Cys	atg Met	tcc Ser	atc Ile 525	atg Met	gcc Ala	ttg Leu	ctg Leu	ctg Leu 530		1882
ctg Leu	ctg Leu	ctc Leu	ctg Leu	ctg Leu 535	cta Leu	ttg Leu	tac Tyr	aag Lys	tat Tyr 540	aag Lys	cag Gln	aag Lys	ccc Pro	aag Lys 545	tac Tyr		1930
Gln	Val	Arg	Trp 550	Lys	Ile	atc Ile	Glu	Ser 555	Tyr	Glu	Gly	Asn	Ser 560	Tyr	Thr	:	1978
ttc Phe	atc Ile	gac Asp 565	ccc Pro	acg Thr	cag Gln	ctg Leu	cct Pro 570	tac Tyr	aac Asn	gag Glu	aag Lys	tgg Trp 575	gag Glu	ttc Phe	ccc Pro	:	2026
cgg Arg	aac Asn 580	aac Asn	ctg Leu	cag Gln	ttt Phe	ggt Gly 585	aag Lys	acc Thr	ctc Leu	gga Gly	gct Ala 590	gga Gly	gcc Ala	ttt Phe	Gly 999	:	2074
aag Lys 595	gtg Val	gtg Val	gag Glu	gcc Ala	acg Thr 600	gcc Ala	ttt Phe	ggt Gly	ctg Leu	ggc Gly 605	aag Lys	gag Glu	gat Asp	gct Ala	gtc Val 610	:	2122
ctg Leu	aag Lys	gtg Val	gct Ala	gtg Val 615	aag Lys	atg Met	ctg Leu	aag Lys	tcc Ser 620	acg Thr	gcc Ala	cat His	gct Ala	gat Asp 625	gag Glu	:	2170
aag	gag	gcc	ctc	atg	tcc	gag	ctg	aag	atc	atg	agc	cac	ctg	ggc	cag	:	2218

Lys	Glu	Ala	Leu 630	Met	Ser	Glu	Leu	Lys 635	Ile	Met	Ser	His	Leu 640	Gly	Gln	
					aac Asn											2266
					gag Glu											2314
					gag Glu 680											2362
					ggc Gly											2410
aaa Lys	tat Tyr	gtc Val	cgc Arg 710	agg Arg	gac Asp	agt Ser	ggc Gly	ttc Phe 715	tcc Ser	agc Ser	cag Gln	ggt Gly	gtg Val 720	gac Asp	acc Thr	2458
					cct Pro											2506
					aag Lys											2554
					agc Ser 760											2602
					cac His											2650
					gcc Ala											2698
					aac Asn											2746
					cca Pro											2794
					tcc Ser 840											2842
					tac Tyr											2890
					gga Gly											2938

aag aat ata tac agc atc atg cag gcc tgc tgg gcc ttg gag ccc acc Lys Asn Ile Tyr Ser Ile Met Gln Ala Cys Trp Ala Leu Glu Pro Thr 885 890 895	2986
cac aga ccc acc ttc cag cag atc tgc tcc ttc ctt cag gag cag gcc His Arg Pro Thr Phe Gln Gln Ile Cys Ser Phe Leu Gln Glu Gln Ala 900 905 910	3034
caa gag gac agg aga gag cgg gac tat acc aat ctg ccg agc agc agc Gln Glu Asp Arg Glu Arg Asp Tyr Thr Asn Leu Pro Ser Ser Ser 915 920 925 930	3082
aga agc ggt ggc agc ggc agc agc agt gag ctg gag gag gag agc Arg Ser Gly Gly Ser Gly Ser Ser Ser Glu Leu Glu Glu Ger 935 940 945	3130
tct agt gag cac ctg acc tgc tgc gag caa ggg gat atc gcc cag ccc Ser Ser Glu His Leu Thr Cys Cys Glu Gln Gly Asp Ile Ala Gln Pro 950 955 960	3178
ttg ctg cag ccc aac aac tat cag ttc tgc tga ggagttgacg acagggagta Leu Leu Gln Pro Asn Asn Tyr Gln Phe Cys 965 970	3231
ceactetece etectecaaa etteaaetee teeatggatg gggcgacaeg gggagaacat	3291
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tgctcacaga ccacactcag ctcaggcccc ttggagcagg atggctcctc taagaatctc	3711
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gcaggaagca caaactcccc caagetgact catcctaact aacagtcacg ccgtgggatg	3951
tctctgtcca cattaaacta acagcattaa tgca	3985

Met Gly Pro Gly Val Leu Leu Leu Leu Val Ala Thr Ala Trp His

Gly Gln Gly Ile Pro Val Ile Glu Pro Ser Val Pro Glu Leu Val Val

<sup>&</sup>lt;210> 8 <211> 972 <212> PRT <213> Homo sapiens

<sup>&</sup>lt;400> 8

20 25 30

Lys Pro Gly Ala Thr Val Thr Leu Arg Cys Val Gly Asn Gly Ser Val 35 40 45

Glu Trp Asp Gly Pro Pro Ser Pro His Trp Thr Leu Tyr Ser Asp Gly 50 55 60

Ser Ser Ser Ile Leu Ser Thr Asn Asn Ala Thr Phe Gln Asn Thr Gly 65 70 75 80

Thr Tyr Arg Cys Thr Glu Pro Gly Asp Pro Leu Gly Gly Ser Ala Ala 85 90 95

Ile His Leu Tyr Val Lys Asp Pro Ala Arg Pro Trp Asn Val Leu Ala
100 105 110

Gln Glu Val Val Phe Glu Asp Gln Asp Ala Leu Leu Pro Cys Leu 115 120 125

Leu Thr Asp Pro Val Leu Glu Ala Gly Val Ser Leu Val Arg Val Arg 130 135 140

Gly Arg Pro Leu Met Arg His Thr Asn Tyr Ser Phe Ser Pro Trp His 145 150 155 160

Gly Phe Thr Ile His Arg Ala Lys Phe Ile Gln Ser Gln Asp Tyr Gln 165 170 175

Cys Ser Ala Leu Met Gly Gly Arg Lys Val Met Ser Ile Ser Ile Arg 180 185 190

Leu Lys Val Gln Lys Val Ile Pro Gly Pro Pro Ala Leu Thr Leu Val

Pro Ala Glu Leu Val Arg Ile Arg Gly Glu Ala Ala Gln Ile Val Cys 210 220

Ser Ala Ser Ser Val Asp Val Asn Phe Asp Val Phe Leu Gln His Asn 225 230 235 240

Asn Thr Lys Leu Ala Ile Pro Gln Gln Ser Asp Phe His Asn Asn Arg 245 250 255

Tyr Gln Lys Val Leu Thr Leu Asn Leu Asp Gln Val Asp Phe Gln His 260 265 270

Ala Gly Asn Tyr Ser Cys Val Ala Ser Asn Val Gln Gly Lys His Ser 275 280 285

- Thr Ser Met Phe Phe Arg Val Val Glu Ser Ala Tyr Leu Asn Leu Ser 290 295 300
- Ser Glu Gln Asn Leu Ile Gln Glu Val Thr Val Gly Glu Gly Leu Asn 305 310 315 320
- Leu Lys Val Met Val Glu Ala Tyr Pro Gly Leu Gln Gly Phe Asn Trp 325 330 335
- Thr Tyr Leu Gly Pro Phe Ser Asp His Gln Pro Glu Pro Lys Leu Ala 340 345 350
- Asn Ala Thr Thr Lys Asp Thr Tyr Arg His Thr Phe Thr Leu Ser Leu 355 360 365 .
- Pro Arg Leu Lys Pro Ser Glu Ala Gly Arg Tyr Ser Phe Leu Ala Arg 370 375 380
- Asn Pro Gly Gly Trp Arg Ala Leu Thr Phe Glu Leu Thr Leu Arg Tyr 385 390 395 400
- Pro Pro Glu Val Ser Val Ile Trp Thr Phe Ile Asn Gly Ser Gly Thr 405 410 415
- Leu Leu Cys Ala Ala Ser Gly Tyr Pro Gln Pro Asn Val Thr Trp Leu 420 425 430
- Gln Cys Ser Gly His Thr Asp Arg Cys Asp Glu Ala Gln Val Leu Gln 435 440 445
- Val Trp Asp Asp Pro Tyr Pro Glu Val Leu Ser Gln Glu Pro Phe His 450 455 460
- Lys Val Thr Val Gln Ser Leu Leu Thr Val Glu Thr Leu Glu His Asn 465 470 475 480
- Gln Thr Tyr Glu Cys Arg Ala His Asn Ser Val Gly Ser Gly Ser Trp 485 490 495
- Ala Phe Ile Pro Ile Ser Ala Gly Ala His Thr His Pro Pro Asp Glu 500 505 510
- Phe Leu Phe Thr Pro Val Val Val Ala Cys Met Ser Ile Met Ala Leu 515 520 525

Leu Leu Leu Leu Leu Leu Leu Leu Tyr Lys Tyr Lys Gln Lys Pro 530 540

Lys Tyr Gln Val Arg Trp Lys Ile Ile Glu Ser Tyr Glu Gly Asn Ser 545 550 555 560

Tyr Thr Phe Ile Asp Pro Thr Gln Leu Pro Tyr Asn Glu Lys Trp Glu 565 570 575

Phe Pro Arg Asn Asn Leu Gln Phe Gly Lys Thr Leu Gly Ala Gly Ala 580 585 590

Phe Gly Lys Val Val Glu Ala Thr Ala Phe Gly Leu Gly Lys Glu Asp 595 600 605

Ala Val Leu Lys Val Ala Val Lys Met Leu Lys Ser Thr Ala His Ala 610 620

Asp Glu Lys Glu Ala Leu Met Ser Glu Leu Lys Ile Met Ser His Leu 625 630 635 640

Gly Gln His Glu Asn Ile Val Asn Leu Leu Gly Ala Cys Thr His Gly 645 650 . 655

Gly Pro Val Leu Val Ile Thr Glu Tyr Cys Cys Tyr Gly Asp Leu Leu 660 : 665 670

Asn Phe Leu Arg Arg Lys Ala Glu Ala Met Leu Gly Pro Ser Leu Ser 675 680 685

Pro Gly Gln Asp Pro Glu Gly Gly Val Asp Tyr Lys Asn Ile His Leu 690 700

Glu Lys Lys Tyr Val Arg Arg Asp Ser Gly Phe Ser Ser Gln Gly Val
705 710 715 720

Asp Thr Tyr Val Glu Met Arg Pro Val Ser Thr Ser Ser Asn Asp Ser 725 730 735

Phe Ser Glu Gln Asp Leu Asp Lys Glu Asp Gly Arg Pro Leu Glu Leu 740 745 750

Arg Asp Leu Leu His Phe Ser Ser Gln Val Ala Gln Gly Met Ala Phe 755 760 765

Leu Ala Ser Lys Asn Cys Ile His Arg Asp Val Ala Ala Arg Asn Val 770 785 780

Leu Leu Thr Asn Gly His Val Ala Lys Ile Gly Asp Phe Gly Leu Ala 785 790 795 800

Arg Asp Ile Met Asn Asp Ser Asn Tyr Ile Val Lys Gly Asn Ala Arg 805 810 815

Leu Pro Val Lys Trp Met Ala Pro Glu Ser Ile Phe Asp Cys Val Tyr 820 825 830

Thr Val Gln Ser Asp Val Trp Ser Tyr Gly Ile Leu Leu Trp Glu Ile 835 840 845

Phe Ser Leu Gly Leu Asn Pro Tyr Pro Gly Ile Leu Val Asn Ser Lys
850 855 860

Phe Tyr Lys Leu Val Lys Asp Gly Tyr Gln Met Ala Gln Pro Ala Phe 865 870 875 880

Ala Pro Lys Asn Ile Tyr Ser Ile Met Gln Ala Cys Trp Ala Leu Glu 885 890 . 895

Pro Thr His Arg Pro Thr Phe Gln Gln Ile Cys Ser Phe Leu Gln Glu . 900 905 910

Gln Ala Gln Glu Asp Arg Arg Glu Arg Asp Tyr Thr Asn Leu Pro Ser 915 920 925

Ser Ser Arg Ser Gly Gly Ser Gly Ser Ser Ser Glu Leu Glu Glu 930 935 940

Glu Ser Ser Ser Glu His Leu Thr Cys Cys Glu Gln Gly Asp Ile Ala 945 950 955 960

Gln Pro Leu Gln Pro Asn Asn Tyr Gln Phe Cys 965 970